

In the Claims:

1-123. (Canceled).

- ~~124~~¹. (Previously presented) An isolated nucleic acid comprising:
- (a) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 326;
 - (b) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 326, lacking its associated signal peptide;
 - (c) the nucleic acid sequence of SEQ ID NO: 325;
 - (d) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 325; or
 - (e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203129;

wherein, said nucleic acid is amplified in colon tumors.

125. (Previously presented) The isolated nucleic acid of Claim 124 comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO:326.

126. (Previously presented) The isolated nucleic acid of Claim 124 comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO:326, lacking its associated signal peptide.

127-128. Canceled.

~~129~~². (Previously presented) The isolated nucleic acid of Claim ~~124~~¹ comprising the nucleic acid sequence of SEQ ID NO: 325.

~~130~~³. (Previously presented) The isolated nucleic acid of Claim ~~124~~¹ comprising the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 325.

~~131~~⁴. (Previously presented) The isolated nucleic acid of Claim ~~124~~¹ comprising the full-length coding sequence of the cDNA deposited under ATCC accession number 203129.

132-134. Canceled

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~~135.~~ (Previously presented) A vector comprising the nucleic acid of Claim ~~124~~¹.

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~~136.~~ (Previously presented) The vector of Claim ~~135~~⁵, wherein said nucleic acid is operably linked to control sequences recognized by a host cell transformed with the vector.

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~~137.~~ (Previously presented) A host cell comprising the vector of Claim ~~135~~⁵.

⁸
~~138.~~ (Previously presented) The host cell of Claim ~~137~~⁷, wherein said cell is a CHO cell, an *E. coli* or a yeast cell.

⁹
~~139.~~ (Currently amended) An isolated nucleic acid molecule consisting of at least a 20 nucleotide fragment of the nucleic acid sequence of SEQ ID NO: 325 or a complement thereof nucleotides in length that specifically hybridizes under stringent conditions to:

- (a) the nucleic acid sequence of SEQ ID NO: ~~325~~ ~~326~~ or a complement thereof; or
- (b) the full-length coding sequence of the cDNA deposited under ATCC accession number 203129 or a complement thereof;

wherein, said stringent conditions use 50% formamide, 5X SSC, 50 mM sodium phosphate (pH 6.8), 0.1% sodium pyrophosphate, 5X Denhardt's solution, sonicated salmon sperm DNA (50 µg/ml), 0.1% SDS, and 10% dextran sulfate at 42°C, and washes at 42°C in 0.2X SSC, at 55°C in 50% formamide followed by a high-stringency wash at 55°C in 0.1X SSC, EDTA; and wherein said isolated nucleic acid molecule is suitable for use as a PCR primer or probe.

¹⁰
~~140.~~ (Currently amended) The isolated nucleic acid molecule of Claim ~~139~~⁹ that is at least 25 nucleotides or greater ~~above~~ in length.

¹¹
~~141~~. (Currently amended) The isolated nucleic acid molecule of Claim ~~139~~⁹ that is at least 30 nucleotides or greater ~~above~~ in length.

¹²
~~142~~. (Currently amended) The isolated nucleic acid molecule of Claim ~~139~~⁹ that is at least 35 nucleotides or greater ~~above~~ in length.